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<110> Burris, Thomas P.

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<211> 2892
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<400> 7

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<400> 8

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Ala Ser Arg Trp Arg Arg Arg Arg Pro Trp Leu Asp Pro Ala Ala Ala
35 40 45

Ala Ala Ala Ala Val Ala Gly Gly Glu Gln Gln Thr Pro Glu Pro Glu
50 55 60

Pro Gly Glu Ala Gly Arg Asp Gly Met Gly Asp Ser Gly Arg Asp Ser
65 70 75 80

```
Arg Ser Pro Asp Ser Ser Ser Pro Asn Pro Leu Pro Gln Gly Val Pro
                85                  90                  95

Pro Pro Ser Pro Pro Gly Pro Pro Leu Pro Pro Ser Thr Ala Pro Ser
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Leu Gly Gly Ser Gly Ala Pro Pro Pro Pro Met Pro Pro Pro Pro
        115                 120                 125

Leu Gly Ser Pro Phe Pro Val Ile Ser Ser Ser Met Gly Ser Pro Gly
    130                 135                 140

Leu Pro Pro Pro Ala Pro Pro Gly Phe Ser Gly Pro Val Ser Ser Pro
145                 150                 155                 160

Gln Ile Asn Ser Thr Val Ser Leu Pro Gly Gly Gly Ser Gly Pro Pro
                165                 170                 175

Glu Asp Val Lys Pro Pro Val Leu Gly Val Arg Gly Leu His Cys Pro
            180                 185                 190

Pro Pro Pro Gly Gly Pro Gly Ala Gly Lys Arg Leu Cys Ala Ile Cys
        195                 200                 205

Gly Asp Arg Ser Ser Gly Lys His Tyr Gly Val Tyr Ser Cys Glu Gly
    210                 215                 220

Cys Lys Gly Phe Phe Lys Arg Thr Ile Arg Lys Asp Leu Thr Tyr Ser
225                 230                 235                 240

Cys Arg Asp Asn Lys Asp Cys Thr Val Asp Lys Arg Gln Arg Asn Arg
                245                 250                 255

Cys Gln Tyr Cys Arg Tyr Gln Lys Cys Leu Ala Thr Gly Met Lys Arg
            260                 265                 270

Glu Ala Val Gln Glu Glu Arg Gln Arg Gly Lys Asp Lys Asp Gly Asp
        275                 280                 285

Gly Glu Gly Ala Gly Gly Ala Pro Glu Glu Met Pro Val Asp Arg Ile
    290                 295                 300

Leu Glu Ala Glu Leu Ala Val Glu Gln Lys Ser Asp Gln Gly Val Glu
305                 310                 315                 320

Gly Pro Gly Gly Thr Gly Gly Ser Gly Ser Ser Pro Asn Asp Pro Val
                325                 330                 335
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Thr Asn Ile Cys Gln Ala Ala Asp Lys Gln Leu Phe Thr Leu Val Glu
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Trp Ala Lys Arg Ile Pro His Phe Ser Ser Leu Pro Leu Asp Asp Gln
355 360 365

Val Ile Leu Leu Arg Ala Gly Trp Asn Glu Leu Leu Ile Ala Ser Phe
370 375 380

Ser His Arg Ser Ile Asp Val Arg Asp Gly Ile Leu Leu Ala Thr Gly
385 390 395 400

Leu His Val His Arg Asn Ser Ala His Ser Ala Gly Val Gly Ala Ile
405 410 415

Phe Asp Arg Val Leu Thr Glu Leu Val Ser Lys Met Arg Asp Met Arg
420 425 430

Met Asp Lys Thr Glu Leu Gly Cys Leu Arg Ala Ile Ile Leu Phe Asn
435 440 445

Pro Asp Ala Lys Gly Leu Ser Asn Pro Ser Glu Val Glu Val Leu Arg
450 455 460

Glu Lys Val Tyr Ala Ser Leu Glu Thr Tyr Cys Lys Gln Lys Tyr Pro
465 470 475 480

Glu Gln Gln Gly Arg Phe Ala Lys Leu Leu Leu Arg Leu Pro Ala Leu
485 490 495

Arg Ser Ile Gly Leu Lys Cys Leu Glu His Leu Phe Phe Phe Lys Leu
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Ile Gly Asp Thr Pro Ile Asp Thr Phe Leu Met Glu Met Leu Glu Ala
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Pro His Gln Leu Ala
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<212> DNA
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<210> 10
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Leu Ser Thr Gly Lys Pro Met Asp Ser His Pro Ser Tyr Thr Asp Thr
        35                  40                  45

Pro Val Ser Ala Pro Arg Thr Leu Ser Ala Val Gly Thr Pro Leu Asn
    50                  55                  60

Ala Leu Gly Ser Pro Tyr Arg Val Ile Thr Ser Ala Met Gly Pro Pro
65                  70                  75                  80

Ser Gly Ala Leu Ala Ala Pro Pro Gly Ile Asn Leu Val Ala Pro Pro
                85                  90                  95

Ser Ser Gln Leu Asn Val Val Asn Ser Val Ser Ser Ser Glu Asp Ile
            100                 105                 110

Lys Pro Leu Pro Gly Leu Pro Gly Ile Gly Asn Met Asn Tyr Pro Ser
        115                 120                 125

Thr Ser Pro Gly Ser Leu Val Lys His Ile Cys Ala Ile Cys Gly Asp
    130                 135                 140

Arg Ser Ser Gly Lys His Tyr Gly Val Tyr Ser Cys Glu Gly Cys Lys
145                 150                 155                 160

Gly Phe Phe Lys Arg Thr Ile Arg Lys Asp Leu Ile Tyr Thr Cys Arg
                165                 170                 175

Asp Asn Lys Asp Cys Leu Ile Asp Lys Arg Gln Arg Asn Arg Cys Gln
            180                 185                 190

Tyr Cys Arg Tyr Gln Lys Cys Leu Val Met Gly Met Lys Arg Glu Ala
        195                 200                 205

Val Gln Glu Glu Arg Gln Arg Ser Arg Glu Arg Ala Glu Ser Glu Ala
    210                 215                 220

Glu Cys Ala Thr Ser Gly His Glu Asp Met Pro Val Glu Arg Ile Leu
225                 230                 235                 240

Glu Ala Glu Leu Ala Val Glu Pro Lys Thr Glu Ser Tyr Gly Asp Met
                245                 250                 255

Asn Met Glu Asn Ser Thr Asn Asp Pro Val Thr Asn Ile Cys His Ala
            260                 265                 270
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Ala Asp Lys Gln Leu Phe Thr Leu Val Glu Trp Ala Lys Arg Ile Pro
        275                 280                 285

His Phe Ser Asp Leu Thr Leu Glu Asp Gln Val Ile Leu Leu Arg Ala
    290                 295                 300

Gly Trp Asn Glu Leu Leu Ile Ala Ser Phe Ser His Arg Ser Val Ser
305                 310                 315                 320

Val Gln Asp Gly Ile Leu Leu Ala Thr Gly Leu His Val His Arg Ser
                325                 330                 335

Ser Ala His Ser Ala Gly Val Gly Ser Ile Phe Asp Arg Val Leu Thr
            340                 345                 350

Glu Leu Val Ser Lys Met Lys Asp Met Gln Met Asp Lys Ser Glu Leu
        355                 360                 365

Gly Cys Leu Arg Ala Ile Val Leu Phe Asn Pro Asp Ala Lys Gly Leu
    370                 375                 380

Ser Asn Pro Ser Glu Val Glu Thr Leu Arg Glu Lys Val Tyr Ala Thr
385                 390                 395                 400

Leu Glu Ala Tyr Thr Lys Gln Lys Tyr Pro Glu Gln Pro Gly Arg Phe
                405                 410                 415

Ala Lys Leu Leu Leu Arg Leu Pro Ala Leu Arg Ser Ile Gly Leu Lys
            420                 425                 430

Cys Leu Glu His Leu Phe Phe Phe Lys Leu Ile Gly Asp Thr Pro Ile
        435                 440                 445

Asp Thr Phe Leu Met Glu Met Leu Glu Thr Pro Leu Gln Ile Thr
    450                 455                 460
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